

AACTAGATGCAGCACCACAATCACTACCAGGTACCAATCATATACCAATAATGTACTAATAATGTACCAATAACTATGGTTTATAAAGATGGTGCATTTAAATCAATATTAGTTCCTTATATTA 125
M V S F K S I L V P Y I

CACTCTTTTAAATGAGCGGTGCTGCTTTGCAAGTGATACCGATCCCGAAGCTGGTGGGCTAGTGAAGCTGGTGGGCTAGTGGAACTGTTGGGCCAGTGAAGCTGGTGGGCTAGTGAAGCT 250
Repeat Sequences
T L F L M S G A V F A S D T D P E A G G P S E A G G P S G T V G P S E A G G P S E A

GGTGGGCTAGTGGAACTGGTGGGCTAGTGAAGCTGGTGGGCTAGTGAAGCTGGTGGGCTAGTGAAGCTGGTGGGCTAGTGAAGCTGGTGGGCTAGTGGAACTGGTGGGCTAGTGAAGCT 375
Repeat Sequences
G G P S G T G W P S E A G G P S E A G G P S E A G G P S E A G G P S G T G W P S G T

TGGTGGGCTAGTGAAGCTGGTGGTGTAGTGAAGCTGGTGGGCTAGTGAAGCTGGTGGGCTAGTGAAGCTGGTGGGCTAGTGAAGCTGGTGGGCTAGTGAAGCTGGTGGGCTAGTGAAGCT 500
Repeat Sequences
G W P S E A G W S S E R F G Y Q L L P Y S R R I V I F N E V C L S Y I Y K H S V M

TATTGGAACGAGATAGGTGAACGATGGTCATAAGACTACATTGAAGAAAAACCAAGGAGAAGAATAATTGAAAAAGAAATGGAAAAATGTTTCTGAAACAATATTCCTTATGAAGAAA 625
I L E R D R V N D G H K D Y I E E K T K E K N K L K K E L E K C F P E Q Y S L M K K

GAAGAATTGGCTAGAATATTGATAATGCATCCACTATCTCTTCAAAATATAAGTTATTGGTTGATGAATATCAACAAGGCCTATGGTACATTGGAAGGTCCAGCTGCTGATAATTTTGACCA 750
E E L A R I F D N A S T I S S K Y K L L V D E I S N K A Y G T L E G P A A D N F D H

TTTCGTAATATATGAAGTCTATTGTACTTAAAGATATGTTATATATTGTGACTTATTATTACAACATTTAATCTATAAATTCTATTATGACAATACCGTTAATGATATCAAGAAAAATTTTG 875
F R N I W K S I V L K D M F I Y C D L L L Q H L I Y K F Y Y D N T V N D I K K N F

ACGAATCCAAATCTAAAGCTTTAGTTTGGGGATAAGATCACTAAAAAGGATGCAGATTATAACACTCATTTTGAGGACATGATTAGGAGTTGAATAGTGCAGCAGAAGAATTTAATAAATT 1000
D E S K S K A L V L R D K I T K D G D Y N T H F E D M I K E L N S A A E E F N K I

GTTGACATCATGATTCCAACTTTGGGATTATGATGAGTATGACAGATTGCAAGTTTCAAACTTTCTTTCAATGATCACCAGAAATCACTAAATCACCAGGTTTCTAATGTAATAATTC 1125
V D I M I S N I G D Y D E Y D S I A S F K P F L S M I T E I T K I T K V S N V I I P

TGGAATTAAGGCACCTAAGTTTAAACGTTTTTTTAAATTTTATTACAAAATAGATGTAATACCAGATGTACATTATTATATTACAAAATTTACACATTATTATGTATGAACGAACGAACAT 1250
G I K A L T L T V F L I F I T K

Fig. 1A

<u>CTCAGTCTTAATGAAGAAATTGGGATAAATATGGAATAGATAAAGTAACATGAGAAAGATGAATATAATATTAGAATATGAAATTTACAGAAATAAATGAAGTAAAGAGTGTATTTGT</u>	1375
<u>AATAATTATAATAAATTAGTATACATGATTATATTACAGATGACTATTGATTATTGTATCAATTAATATTGATTATTAATGATATCATATGTATATGTTAATGATTGATTGTTATACGT</u>	1500
<u>TGTGAATATGTTATATAATGACACTATAATAATTAATATAATGTAGAGGATATTTTTTAAATAGTATTAAATGAATATTATAGTTATAATTATAATAATGTAGATAAAATGACATTAAATT</u>	1625
<u>GAATGTTAAATTGAATGTATGTAAAAATATGATTATAATCTGAATTGATTAATAATATAATATTCTACAATTAATTATTTTGTAAATTATAAATGATTATTAATCTTTGAATTATT</u>	1750
<u>ATAAATAATTATACCTCATTAAATATTTACATAAATTTCCAAATTATTATCCTTTATCTTAATGTTATCCAATTTACACATCTTCTTCATTACAATATTTTTACTAATCCTGTATGC</u>	1875
<u>TCATATTCATATTCCTTAGAAATATAACGAAATAGATGTAACCTCGCCACTTACAAGTAACTACCATCAATATAATAATGAATACCATTGATCGGTATATCTTTATATTTTATC</u>	2000
<u>ATATTTTATTTGTGATTATTCATTCAATTTGTATCATTATCAATGAGAGAAATAATAGCAGAAAGATCCTTCTATAGAAACATAAAATTCATTAATACTGGATTATTATGTTTGAAGTATA</u>	2125
<u>GATGTTAAATCAATAACACTACCAGTTGGTAATTAGCATTGTCATCAATTCATTAATATAATCAGAAATTTGATTTTATCAATTTTATTCGGATGTGATAATTTATTTGTTCTGATTCAT</u>	2250
<u>CGATCATGTATACAAATACTATTGTTAAAGTTCCCTATCCTTATAATTAAGTGGCCAAATAAGATTGGCATTAAATACATTAGTAGTGTGTATTGTAATAGTATCATTAGTGGTACTGACA</u>	2375
<u>GTTGTTATAGGTTTTGATTCCATAATGAACATCATTTTATCTACACAATACA</u>	2430

Fig. 1B

Figure 2A shows the results of the ELISA assay for the detection of anti-Babesia antibodies in serum samples. The y-axis represents the optical density (OD) at 450 nm, ranging from 0 to 3. The x-axis represents the serum ID, categorized into Babesia and Normals. The legend indicates four different antigen preparations: BABS-1 (diagonal lines), BMNI-3 (white), BABS-4 (diagonal lines), and BMNI-6 (checkered). The results show that all Babesia samples (DHTY, HGH, #252, #145, CORIXA 1, CORIXA 2, CORIXA 4, CORIXA 5, CORIXA 6, AT4-0043, AT4-0052, AT4-0053, AT4-0062, AT4-0070) have OD values significantly higher than the Normals, indicating a positive result for anti-Babesia antibodies. The OD values for Babesia samples are generally higher than those for Normals, with some samples showing OD values above 2.5.

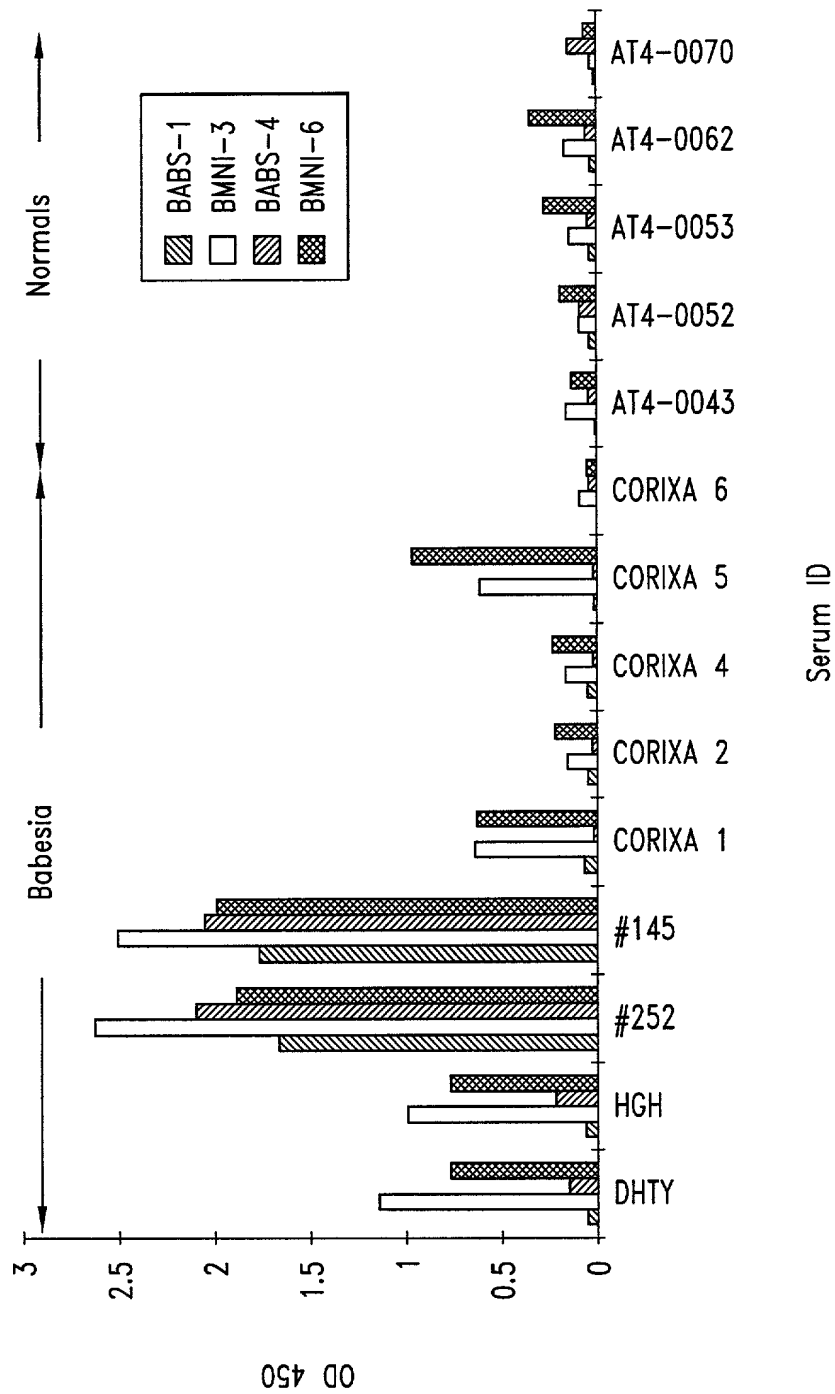


Fig. 2A



Fig. 2B

BMNI-1 BMNI-2 BMNI-3 BMNI-4 BMNI-5 BMNI-6 BMNI-7 BMNI-8 BMNI-9 BMNI-10 BMNI-11 BMNI-12 BMNI-13 BMNI-14 BMNI-15

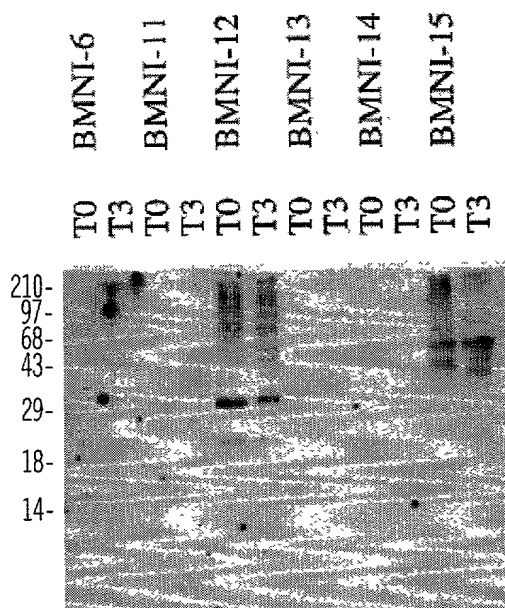
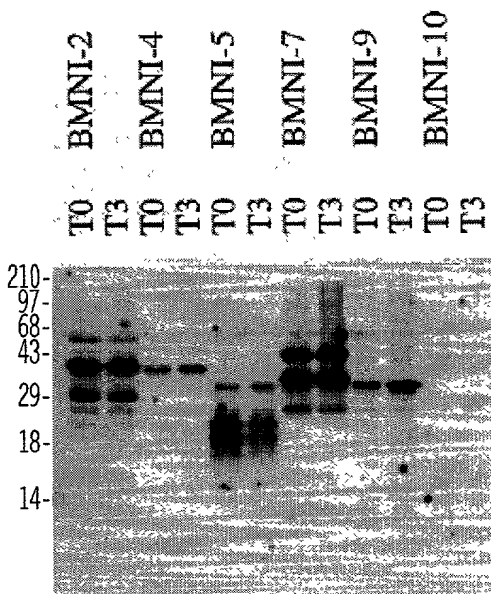
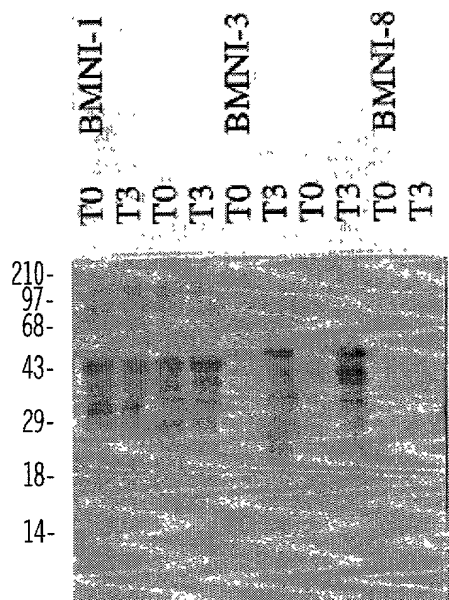


Fig. 4

1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20

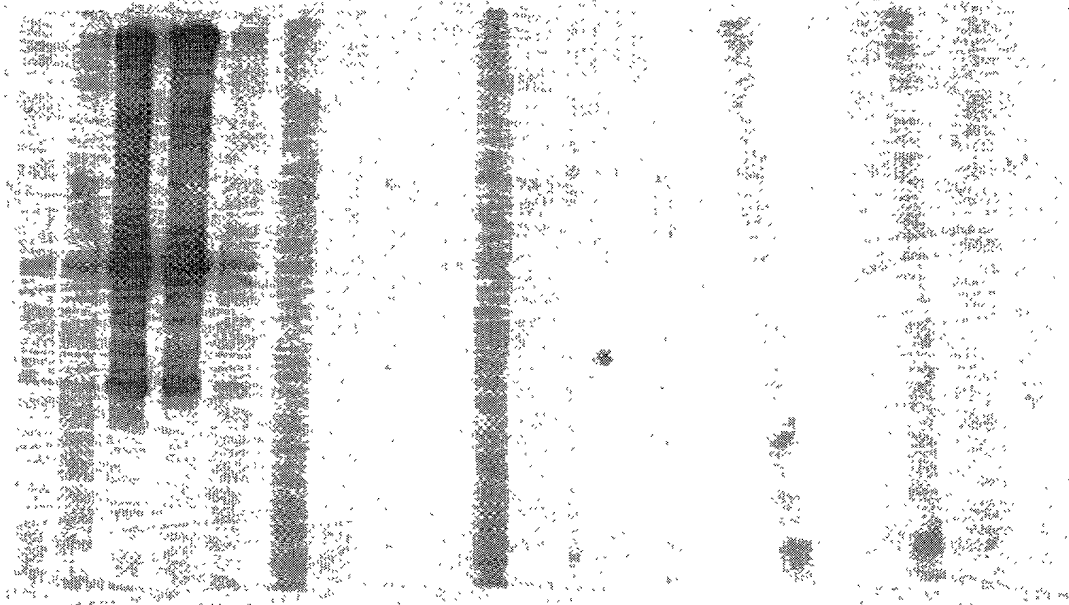


Fig. 5

BI254AGDTDREA	GGPSGTVGP.
BI1053GDTDREA	GGPSGTVGP.
BI2227AGDTDREA	GGPSGTVGP.SEAGGPSEA
BI2259AGDTDREA	GGPSGTVGP.SEAGGPSEA
BI2253EA	GGPSGTVGP.SEAGGPSEA
GRAC,SGDTDREA	GGPSGTVGP.SEAGG PSEAGGPSEA
FISH,SAGDTDREA	GGPSGTVGPS	SAGGPSEAGG	PSEAGGPSEA
MN1HAMAGDTDREA	GGPSGTVGP.SEA
MN2AGDTDREA	GGPSGTVGP.
MN1PATAGDTDREA	GGPSGTVGP.SEA
Bmni-6	YITLFLMSG	VFAGDTDREA	GGPSGTVGP.SEA
MN3AGDTDREA	GGPSGTVGP.SEAGGPSEA
MR.TAGDTDREA	GGPSGTVGP.SEAGGPSEA
	51				100
BI254	...	SEAGGPS	EAGGPSGTVG	PSEAGGPSEA	GGPSGTGWPS EAGGPSGTVG
BI1053	...	SEAGGPS	EAGGPSGTVG	PSEAGGPSEA	GGPSGTGWPS EAGGPSGTVG
BI2227	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS	EAGGPSEAGW
BI2259	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS	EAGGPSEAGW
BI2253	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS	EAGGPSEAGW
GRAC,S	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS	EAGGPSEAGW
FISH,S	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS	EAGGPSEAGW
MN1HAM	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS	EAGGPSGTGW
MN2	...	SEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS EAGGPSGTGW
MN1PAT	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS	EAGGPSGTGW
Bmni-6	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSHAGGPS	EAGGPSGTGW
MN3	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS	EAGGPSGTGW
MR.T	GGPSEAGGPS	EAGGPSEAGG	PSEAGGPSEA	GGPSEAGGPS	EAGGPSGTGW
	101				150
BI254	PSEAGGP...S	EAGGPSGTGW	PSGTGWPEV	GWPSERFGYQ
BI1053	PSEAGGP...S	EAGGPSGTGW	PSGTGWPEV	GWPSERFGYQ
BI2227	PSEAGWPSEA	GGPSGTGWPS	EAGWPSEAGW	PSEAGWPSEA	GW.....
BI2259	PSEAGWPSEA	GGPSGTGWPS	EAGWPSEAGW	PSEAGWPSEA	GWPSERFGYQ
BI2253	PSEAGWPSEA	GGPSGTGWPS	EAGWPSEAGW	PSEAGWPSEA	GWPSER....
GRAC,S	PSEAGWPSEA	GGPSGTGWPS	EAGWPSEAGW	PSEAGWPSEA	GWPSERFGYQ
FISH,S	PSEAGWPSEA	GGPSGTGWPS	EAGWPSEAGW	PSEAGWPSEA	GWPSERFGYQ
MN1HAM	PSEAGWP...S	EAGWPSEAGW	PSEAGWPSEA	GWPSERFGYQ
MN2	PSEAGWP...S	EAGWPSEAGW	PSEAGWPSEA	GW.....
MN1PAT	PSEAGWP...S	EAGWPSEAGW	PSEAGWPSEA	GWPSERFGYQ
Bmni-6	PSEAGWP...S	EAGWPSEAGW	PSEAGWPSEA	GWPSERFGYQ
MN3	PSEAGWP...S	EAGWPSEAGW	PSEAGWPSEA	GWPSERFGYQ
MR.T	PSEAGWP...S	EAGWPSEAGW	PSEAGWPSEA	GWPSERFGYQ

Fig. 6A

	151	177
BI254	LLWYSRRIVI	
BI1053	LLWYSRRIVI	
BI2227	
BI2259	LLWYSRRIVI	
BI2253	
GRAC,S	LLWYS.....	
FISH,S	
MN1HAM	LLWYSRRIVI	
MN2	
MN1PAT	LLWYS.....	
Bmni-6	LLWYSRRIVI FNEIYLSHIY EHSVMIL	
MN3	LLWYSR.....	
MR.T	LLWYSR.....	

Fig. 6B